

SEQUENCE LISTING

OIPE 40	Kuja-Panula Kiiltomaki, Rauvala, He	Marjaana					
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Lys Gln Gln Leu Pro Asn Val Pro His Ser Leu Pro Ser Tyr Thr Ala

Leu Leu Asp Leu Ser His Asn Asn Leu Ser Arg Leu Arg Ala Glu Trp 75 70 Thr Pro Thr Arg Leu Thr Gln Leu His Ser Leu Leu Leu Ser His Asn His Leu Asn Phe Ile Ser Ser Glu Ala Phe Ser Pro Val Pro Asn Leu 100 105 Arg Tyr Leu Asp Leu Ser Ser Asn Gln Leu Arg Thr Leu Asp Glu Phe 115 120 Leu Phe Ser Asp Leu Gln Val Leu Glu Val Leu Leu Tyr Asn Asn His Ile Met Ala Val Asp Arg Cys Ala Phe Asp Asp Met Ala Gln Leu 150 155 Gln Lys Leu Tyr Leu Ser Gln Asn Gln Ile Ser Arg Phe Pro Leu Glu 165 170 Leu Val Lys Glu Gly Ala Lys Leu Pro Lys Leu Thr Leu Leu Asp Leu 185 Ser Ser Asn Lys Leu Lys Asn Leu Pro Leu Pro Asp Leu Gln Lys Leu 195 Pro Ala Trp Ile Lys Asn Gly Leu Tyr Leu His Asn Asn Pro Leu Asn 215 Cys Asp Cys Glu Leu Tyr Gln Leu Phe Ser His Trp Gln Tyr Arg Gln 230 235 Leu Ser Ser Val Met Asp Phe Gln Glu Asp Leu Tyr Cys Met Asn Ser Lys Lys Leu His Asn Val Phe Asn Leu Ser Phe Leu Asn Cys Gly Glu 265 Tyr Lys Glu Arg Ala Trp Glu Ala His Leu Gly Asp Thr Leu Ile Ile 275 280 Lys Cys Asp Thr Lys Gln Gln Gly Met Thr Lys Val Trp Val Thr Pro Ser Asn Glu Arg Val Leu Asp Glu Val Thr Asn Gly Thr Val Ser Val 315 Ser Lys Asp Gly Ser Leu Leu Phe Gln Gln Val Gln Val Glu Asp Gly 325 Gly Val Tyr Thr Cys Tyr Ala Met Gly Glu Thr Phe Asn Glu Thr Leu 345 Ser Val Glu Leu Lys Val His Asn Phe Thr Leu His Gly His His Asp

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Ser Met Leu Ser 420	Thr Thr Pro	Asn His Asp 425		ly Gly Asp 30
Lys Asp Asp Gly 435	Phe Asp Arg	Arg Val Ala 440	Phe Leu Glu Pr 445	co Ala Gly
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Asn Leu Phe Arg Leu Ile Lys Arg Leu Asp Leu Ser Tyr Asn Arg Ile

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75

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- Thr Asp Ile Val Ser Cys Thr Asn Lys Asn Leu Ser Lys Val Pro Gly 50 55 60
- Asn Leu Phe Arg Leu Ile Lys Arg Leu Asp Leu Ser Tyr Asn Arg Ile 65 70 75 80
- Gly Leu Leu Asp Ser Glu Trp Ile Pro Val Ser Phe Ala Lys Leu Asn 85 90 95
- Thr Leu Ile Leu Arg His Asn Asn Ile Thr Ser Ile Ser Thr Gly Ser 100 105 110
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- Met His His Ile Asn Leu Val Pro Gly Lys Gln Leu Arg Gly Ile Tyr 210 215 220
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			tca Ser												384
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			cac His												624
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ccg gcc Pro Ala 305	_	-		_		_	_	_	-	-					960
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Leu Leu Asp Leu 115	Ser Ser A	sn Thr Leu 120	Arg Ala L	Leu Gly Arg 125	His Asp
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Pro Gly Gly Leu Gln Leu Lys Ala Gly Ser Glu Ser Ala Ser Ser Ile

490

140

135

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			_	_			_			_	tcc Ser		_			1776
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Pro Ala Trp Val Lys Asn Gly Leu Tyr Leu His Asn Asn Pro Leu Glu

210 215 220

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Ser Arg Trp Gln 580	Gln Gly Asn	Val Phe Ser 585	Cys Ser Val	Met His 590	Glu
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					tac Tyr									1440
					gag Glu									1488
					cac His									1536
					aaa Lys									1584
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Pro Pro Ser 545		gag ctg Glu Leu 550			Asn Gl						1680
ctg gtc aaa Leu Val Lys				Asp 1							1728
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Ser Thr Thr Pro Asn Leu Lys Cys Leu Asp Leu Ser Ser Asn Arg Leu 115 120 Lys Ser Val Lys Ser Ala Thr Phe Gln Glu Leu Lys Ala Leu Glu Val 135 Leu Leu Leu Tyr Asn Asn His Ile Ser Tyr Leu Asp Pro Ala Ala Phe 155 150 Gly Gly Leu Ser His Leu Gln Lys Leu Tyr Leu Ser Gly Asn Phe Leu 165 Thr Gln Phe Pro Met Asp Leu Tyr Thr Gly Arg Phe Lys Leu Ala Asp Leu Thr Phe Leu Asp Val Ser Tyr Asn Arg Ile Pro Ser Ile Pro Met 200 His His Ile Asn Leu Val Pro Gly Arg Gln Leu Arg Gly Ile Tyr Leu His Gly Asn Pro Phe Val Cys Asp Cys Ser Leu Tyr Ser Leu Leu Ile 230 235 Phe Trp Tyr Arg Arg His Phe Ser Ser Val Met Asp Phe Lys Asn Asp 245 250 Tyr Thr Cys Arg Leu Trp Ser Asp Ser Arg His Ser His Gln Leu Gln 265 Leu Leu Gln Glu Ser Phe Leu Asn Cys Ser Tyr Ser Val Ile Asn Gly 280 Ser Phe His Ala Leu Gly Phe Ile His Glu Ala Gln Val Gly Glu Arg 295 Ala Ile Val His Cys Asp Ser Lys Thr Gly Asn Gly Asn Thr Asp Phe 310 315 Ile Trp Val Gly Pro Asp Asn Arg Leu Leu Glu Pro Asp Lys Asp Met 330 325 Gly Asn Phe Arg Val Phe Tyr Asn Gly Ser Leu Val Ile Glu Asn Pro 345 Gly Phe Glu Asp Ala Gly Val Tyr Ser Cys Ile Ala Met Asn Arg Gln Arg Leu Leu Asn Glu Thr Val Asp Ile Met Ile Asn Val Ser Asn Phe Thr Ile Asn Arg Ser His Ala His Glu Ala Ala Ala Asp Pro Ile 390 395 Glu Gly Arg Gly Gly Gly Gly Asp Pro Lys Ser Cys Asp Lys Pro 405 410 415

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His Thr Cys Pro Pro Cys Pro Ala Pro Glu Leu Leu Gly Gly Pro Ser 420 425 430
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Lys Thr Lys Pro Arg Glu Glu Gln Tyr Asn Ser Thr Tyr Arg Val Val 485 490 495

Ser Val Leu Thr Val Leu His Gln Asp Trp Leu Asn Gly Lys Glu Tyr 500 505 510

Lys Cys Lys Val Ser Asn Lys Ala Leu Pro Ala Pro Ile Glu Lys Thr 515 520 525

Ile Ser Lys Ala Lys Gly Gln Pro Arg Glu Pro Gln Val Tyr Thr Leu 530 540

Pro Pro Ser Arg Asp Glu Leu Thr Lys Asn Gln Val Ser Leu Thr Cys 545 550 555 560

Leu Val Lys Gly Phe Tyr Pro Ser Asp Ile Ala Val Glu Trp Glu Ser 565 570 575

Asn Gly Gln Pro Glu Asn Asn Tyr Lys Thr Thr Pro Pro Val Leu Asp 580 585 590

Ser Asp Gly Ser Phe Phe Leu Tyr Ser Lys Leu Thr Val Asp Lys Ser 595 600 605

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	gaa Glu										336
	ctt Leu										384
	gat Asp 130										432
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	ctc Leu										528
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-					gaa Glu											768
			_		cgc Arg					_		_		_		816
-			-	-	gct Ala					_		_				864
		_			cag Gln		_				-			_		912
	_				gcc Ala 310				_	_			_			960
					gat Asp											1008
	_				gtg Val			_		-		-			_	1056
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	-			_	gct Ala	-										1152
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					cac His											1248
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					acc Thr											1344
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Thr Tyr Arq Val Val Ser Val Leu Thr Val Leu His Gln Asp Trp Leu
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Asn Gly Lys Glu Tyr Lys Cys Lys Val Ser Asn Lys Ala Leu Pro Ala
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Pro Ile Glu Lys Thr Ile Ser Lys Ala Lys Gly Gln Pro Arg Glu Pro
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Gln Val Tyr Thr Leu Pro Pro Ser Arg Asp Glu Leu Thr Lys Asn Gln
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300

His Ala Gln Val Gly Gln Ser Leu Arg Leu Phe Cys Asn Thr Ser Val

295

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Ala Ala Asp 385	Pro Ile	Glu Gl 390	y Arg	Gly	Gly	Gly 395	Gly	Gly	Asp	Pro	Lys 400
Ser Cys Asp	Lys Pro 405		r Cys	Pro	Pro 410	Cys	Pro	Ala	Pro	Glu 415	Leu
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Thr Tyr Arg	Val Val 485		l Leu	Thr	Val 490	Leu	His	Gln	Asp	Trp 495	Leu
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Gln Val Tyr 530	Thr Leu	Pro Pr 53		Arg	Asp	Glu	Leu 540	Thr	Lys	Asn	Gln
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Val Glu Trp	Glu Ser 565		y Gln	Pro	Glu 570	Asn	Asn	Tyr	Lys	Thr 575	Thr
Pro Pro Val	Leu Asp 580	Ser As	p Gly	Ser 585	Phe	Phe	Leu	Tyr	Ser 590	Lys	Leu
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Val Met His Glu Ala Leu His Asn His Tyr Thr Gln Lys Ser Leu Ser

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Cys 225	Asp	Cys	Lys	Leu	Tyr 230	Gln	Leu	Phe	Ser	His 235	Trp	Gln	Tyr	Arg	Gln 240
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Gly Phe Glu Asp Ala Gly Val Tyr Ser Cys Ile Ala Met Asn Arg Gln 360 Arg Leu Leu Asn Glu Thr Val Asp Ile Met Ile Asn Val Ser Asn Phe 375 Thr Ile Asn Arg Ser His Ala His Glu Ala Phe Asn Thr Ala Phe Thr 390 395 Thr Leu Ala Ala Cys Val Ala Ser Ile Val Leu Val Leu Leu Tyr Leu 405 410 Tyr Leu Thr Pro Cys Pro Cys Lys Cys Lys Ala Lys Arg Gln Lys Asn 425 Thr Leu Ser Gln Ser Ser Ala His Ser Ser Ile Leu Ser Pro Gly Pro 440 Thr Gly Asp Ala Ser Ala Asp Asp Arg Lys Ala Gly Lys Arg Val Val Phe Leu Glu Pro Leu Lys Asp Thr Ala Ala Gly Gln Asn Gly Lys Val 470 475 Lys Leu Phe Pro Ser Glu Thr Val Ile Ala Glu Gly Ile Leu Lys Ser 485 Thr Arg Ala Lys Ser Asp Ser Asp Ser Val Asn Ser Val Phe Ser Asp 505 Thr Pro Phe Val Ala Ser Thr 515 <210> 17 <211> 1524 <212> DNA <213> Mus musculus <220> <221> CDS <222> (1)..(1524) <400> 17 atg gcc tgg cta gtg cta tca ggt ata cta cta tgc atg ttg ggt gct 48 Met Ala Trp Leu Val Leu Ser Gly Ile Leu Leu Cys Met Leu Gly Ala 10 15 gga ttg ggc act tca gac ttg gag gat gtt ctg cct cct gct ccc cac Gly Leu Gly Thr Ser Asp Leu Glu Asp Val Leu Pro Pro Ala Pro His 20 25

Gly Asn Phe Arg Val Phe Tyr Asn Gly Ser Leu Val Ile Glu Asn Pro

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					gac Asp											192
-		-	_	_	cac His 70		_			_	_		_			240
					cgg Arg											288
					cat His											336
		_	_		tct Ser		_			_					-	384
_	_		-		gag Glu	_		_			_					432
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					ctc Leu											672
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Ser	Glu	Ser	Arg 260	Val	Arg	Phe	Phe	Glu 265	His	Ser	Arg	Val	Phe 270	Lys	Asn	
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	-		-		_		_	tcc Ser		-			_			960
								atc Ile								1008
								cag Gln 345		-		-				1056
_	-	_			_	_		cac His		_						1104
				_		_		gag Glu								1152
								gtg Val								1200
								ggc Gly								1248
								cgg Arg 425								1296
								agc Ser								1344
								cat His								1392
								cag Gln								1440
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1524

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Asn Cys Pro Asp Ile Cys Ile Cys Ala Ala Asp Val Leu Ser Cys Ala 35 40 45

Gly Arg Gly Leu Gln Asp Leu Pro Val Ala Leu Pro Thr Thr Ala Ala 50 55 60

Glu Leu Asp Leu Ser His Asn Ala Leu Lys Arg Leu His Pro Gly Trp 65 70 75 80

Leu Ala Pro Leu Ser Arg Leu Arg Ala Leu His Leu Gly Tyr Asn Lys
85 90 95

Leu Glu Val Leu Gly His Gly Ala Phe Thr Asn Ala Ser Gly Leu Arg 100 105 110

Thr Leu Asp Leu Ser Ser Asn Met Leu Arg Met Leu His Thr His Asp 115 120 125

Leu Asp Gly Leu Glu Glu Leu Glu Lys Leu Leu Phe Asn Asn Ser 130 135 140

Leu Met His Leu Asp Leu Asp Ala Phe Gln Gly Leu Arg Met Leu Ser 145 150 155 160

His Leu Tyr Leu Ser Cys Asn Glu Leu Ser Ser Phe Ser Phe Asn His 165 170 175

Leu His Gly Leu Gly Leu Thr Arg Leu Arg Thr Leu Asp Leu Ser Ser 180 185 190

Asn Trp Leu Lys His Ile Ser Ile Pro Glu Leu Ala Ala Leu Pro Thr 195 200 205

Tyr Leu Lys Asn Arg Leu Tyr Leu His Asn Asn Pro Leu Pro Cys Asp 210 215 220

Cys Ser Leu Tyr His Leu Leu Arg Arg Trp His Gln Arg Gly Leu Ser 225 230 235 240

Ala Leu His Asp Phe Glu Arg Glu Tyr Thr Cys Leu Val Phe Lys Val Ser Glu Ser Arg Val Arg Phe Phe Glu His Ser Arg Val Phe Lys Asn 265 260 Cys Ser Val Ala Ala Pro Gly Leu Glu Leu Pro Glu Glu Gln Leu 280 His Ala Gln Val Gly Gln Ser Leu Arg Leu Phe Cys Asn Thr Ser Val 295 Pro Ala Thr Arg Val Ala Trp Val Ser Pro Lys Asn Glu Leu Leu Val 310 Ala Pro Ala Ser Gln Asp Gly Ser Ile Ala Val Leu Ala Asp Gly Ser Leu Ala Ile Gly Arg Val Gln Glu Gln His Ala Gly Val Phe Val Cys 340 Leu Ala Ser Gly Pro Arg Leu His His Asn Gln Thr Leu Glu Tyr Asn 360 Val Ser Val Gln Lys Ala Arg Pro Glu Pro Glu Thr Phe Asn Thr Gly 375 380 Phe Thr Thr Leu Leu Gly Cys Ile Val Gly Leu Val Leu Leu Leu 390 385 Tyr Leu Phe Ala Pro Pro Cys Arg Gly Cys Cys His Cys Cys Gln Arg 410 Ala Cys Arg Asn Arg Cys Trp Pro Arg Ala Ser Ser Pro Leu Gln Glu 420 425 Leu Ser Ala Gln Ser Ser Met Leu Ser Thr Thr Pro Pro Asp Ala Pro 435 Ser Arg Lys Ala Ser Val His Lys His Val Val Phe Leu Glu Pro Gly 455 Lys Lys Gly Leu Asn Gly Arg Val Gln Leu Ala Val Ala Glu Asp Phe 465 470 Asp Leu Cys Asn Pro Met Gly Leu Gln Leu Lys Ala Gly Ser Glu Ser 490 Ala Ser Ser Thr Gly Ser Glu Gly Leu Val Met Ser 500 505

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		195					200					205				
aaa d Lys I																672
ggc a Gly I 225																720
aca q Thr (768
gaa q Glu <i>A</i>																816
acc a	_		_	_	-								-			864
gcc a Ala 1		_		-	_	_		_								912
ggc t Gly 3 305																960
gac (Asp (_	_	_	_	_	_	-	_			-	_			1008
tgt a Cys <i>I</i>																1056
gct a Ala 1	_								-				-		_	1104
ctc o Leu I																1152
cct o Pro I 385		_	_		_	-	_	-		-			-	_	_	1200
atc a				_	_		-	_			-			_		1248
ctc (Leu l																1296

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							aca Thr						1440
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							gaa Glu						1920
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	ctg Leu															2016
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	gag Glu 690					-		_	_				_		-	2112
	atc Ile															2160
	gcg Ala			_			_						-			2208
	gtt Val															2256
	aaa Lys															2304
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	gtc Val		-			_					_		_			2448
	tgt Cys															2496
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	cat His 850															2592
	gag Glu															2640
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Met .	Ala	Leu	Glu	Ser 885	Ile	Leu	His	Arg	Ile 890	Tyr	Thr	His	Gln	Ser 895	Asp	
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aag Lys																2784
aaa Lys		-	_			_				-			-			2832
atg Met 945																2880
ttc Phe																2928
cgc Arg			_		_		-	_	_	_		_		_		2976
aca Thr						Arg	_	_	_	_	Glu	-	-	-	_	3024
gac Asp					Asp					Pro						3072
agc Ser 1025	Ser			Thr					Leu		_		ctg Leu	Ser	-	3120
acc Thr			Asn					Cys					Gly			3168
agc Ser	-	Pro		_	_	-	Ser		_	_	_	Tyr	_		-	3216
ccc Pro	Thr					Glu					Asp					3264
gtg Val 1		-			Asn	_		_		Lys			-			3312
gtg Val																3360

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Leu Ser Leu Gln Arc	g Met Phe Asn Ası 55	n Cys Glu Val Val 60	Leu Gly Asn
Leu Glu Ile Thr Tyr 65	r Val Gln Arg Ası 70	n Tyr Asp Leu Ser 75	Phe Leu Lys 80
Thr Ile Gln Glu Vai	. Ala Gly Tyr Va	L Leu Ile Ala Leu 90	Asn Thr Val

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Asp Gly Val Arg Lys Cys Lys Cys Glu Gly Pro Cys Arg Lys Val Cys Asn Gly Ile Gly Ile Gly Glu Phe Lys Asp Ser Leu Ser Ile Asn Ala Thr Asn Ile Lys His Phe Lys Asn Cys Thr Ser Ile Ser Gly Asp Leu His Ile Leu Pro Val Ala Phe Arg Gly Asp Ser Phe Thr His Thr Pro Pro Leu Asp Pro Gln Glu Leu Asp Ile Leu Lys Thr Val Lys Glu Ile Thr Gly Phe Leu Leu Ile Gln Ala Trp Pro Glu Asn Arg Thr Asp Leu His Ala Phe Glu Asn Leu Glu Ile Ile Arg Gly Arg Thr Lys Gln His Gly Gln Phe Ser Leu Ala Val Val Ser Leu Asn Ile Thr Ser Leu Gly Leu Arg Ser Leu Lys Glu Ile Ser Asp Gly Asp Val Ile Ile Ser Gly Asn Lys Asn Leu Cys Tyr Ala Asn Thr Ile Asn Trp Lys Lys Leu Phe Gly Thr Ser Gly Gln Lys Thr Lys Ile Ile Ser Asn Arg Gly Glu Asn Ser Cys Lys Ala Thr Gly Gln Val Cys His Ala Leu Cys Ser Pro Glu Gly Cys Trp Gly Pro Glu Pro Arg Asp Cys Val Ser Cys Arg Asn Val Ser Arg Gly Arg Glu Cys Val Asp Lys Cys Asn Leu Leu Glu Gly Glu Pro Arg Glu Phe Val Glu Asn Ser Glu Cys Ile Gln Cys His Pro

Glu Cys Leu Pro Gln Ala Met Asn Ile Thr Cys Thr Gly Arg Gly Pro 565 570 575

Asp Asn Cys Ile Gln Cys Ala His Tyr Ile Asp Gly Pro His Cys Val 580 580 590

Lys Thr Cys Pro Ala Gly Val Met Gly Glu Asn Asn Thr Leu Val Trp 595 600 605

Lys Tyr Ala Asp Ala Gly His Val Cys His Leu Cys His Pro Asn Cys 610 615 620

Thr Tyr Gly Cys Thr Gly Pro Gly Leu Glu Gly Cys Pro Thr Asn Gly 625 630 635 640

Pro Lys Ile Pro Ser Ile Ala Thr Gly Met Val Gly Ala Leu Leu Leu 645 650 655

Leu Leu Val Val Ala Leu Gly Ile Gly Leu Phe Met Arg Arg Arg His 660 665 670

Ile Val Arg Lys Arg Thr Leu Arg Arg Leu Leu Gln Glu Arg Glu Leu 675 680 685

Val Glu Pro Leu Thr Pro Ser Gly Glu Ala Pro Asn Gln Ala Leu Leu 690 695 700

Arg Ile Leu Lys Glu Thr Glu Phe Lys Lys Ile Lys Val Leu Gly Ser 705 710 715 720

Gly Ala Phe Gly Thr Val Tyr Lys Gly Leu Trp Ile Pro Glu Gly Glu 725 730 735

Lys Val Lys Ile Pro Val Ala Ile Lys Glu Leu Arg Glu Ala Thr Ser 740 745 750

Pro Lys Ala Asn Lys Glu Ile Leu Asp Glu Ala Tyr Val Met Ala Ser 755 760 765

Val Asp Asn Pro His Val Cys Arg Leu Leu Gly Ile Cys Leu Thr Ser 770 780

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Trp	Cys	Val	Gln 820	Ile	Ala	Lys	Gly	Met 825	Asn	Tyr	Leu	Glu	Asp 830	Arg	Arg	
Leu	Val	His 835	Arg	Asp	Leu	Ala	Ala 840	Arg	Asn	Val	Leu	Val 845	Lys	Thr	Pro	
Gln	His 850	Val	Lys	Ile	Thr	Asp 855	Phe	Gly	Leu	Ala	Lys 860	Leu	Leu	Gly	Ala	
Glu 865	Glu	Lys	Glu	Tyr	His 870	Ala	Glu	Gly	Gly	Lys 875	Val	Pro	Ile	Lys	Trp 880	
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Thr	Asp	Ser 995	Asn	Phe	Tyr	Arg	Ala 1000		Met	. Asp	Glu	ı Glu 100		sp Me	et Asp	

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Pro Ala Ser Arg Ala Leu Glu Glu Lys Lys Val Cys Gln Gly Thr Ser
aac aag ctc acg cag ttg ggc act ttt gaa gat cat ttt ctc agc ctc
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Asn Lys Leu Thr Gln Leu Gly Thr Phe Glu Asp His Phe Leu Ser Leu
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Gln Arg Met Phe Asn Asn Cys Glu Val Val Leu Gly Asn Leu Glu Ile
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Thr Tyr Val Gln Arg Asn Tyr Asp Leu Ser Phe Leu Lys Thr Ile Gln
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                         90
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Pro Leu Glu Asn Leu Gln Ile Ile Arg Gly Asn Met Tyr Tyr Glu Asn
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Ser Tyr Ala Leu Ala Val Leu Ser Asn Tyr Asp Ala Asn Lys Thr Gly
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	aac Asn								751
	agc Ser								799
	atc Ile								847
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	gcc Ala 310								1135
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	cac His								1279

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					gct Ala											1423
				-	atc Ile 425		_				_				_	1471
			-	_	gtc Val	_	-					_			-	1519
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	_	-		-	aat Asn							_				1615
					aaa Lys			_				-		-	_	1663
					gtc Val 505											1711
					agg Arg											1759
		_	_		gac Asp	_	_			_						1807
					tct Ser											1855
					atc Ile											1903
					tac Tyr 585											1951
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Pro	Ala	Gly	Val	Met 600	Gly	Glu	Asn	Asn	Thr 605	Leu	Val	Trp	Lys	Tyr 610	Ala	
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					ctt Leu	-		_		-				-		2095
			-		ggg Gly	-			-			_	_	_		2143
					ggc Gly 665											2191
					agg Arg	_	_	-								2239
					gaa Glu					_		_			-	2287
_	-		_		aaa Lys	_				_						2335
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					aag Lys 745											2431
					gat Asp											2479
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gac gga atc cct a Asp Gly Ile Pro				
cgc ctc cct cag Arg Leu Pro Gln : 935		-	, ,	-
gtc aag tgc tgg . Val Lys Cys Trp 1 950	Met Ile Asp		-	
ttg atc atc gaa Leu Ile Ile Glu 965				
gtc att cag ggg o Val Ile Gln Gly 2 980				
aac ttc tac cgt of Asn Phe Tyr Arg 1			Asp Met Asp Asp	
gat gcc gac gag Asp Ala Asp Glu ' 1015				
tcc acg tca cgg a Ser Thr Ser Arg ' 1030	Thr Pro Leu			_
aat tcc acc gtg o Asn Ser Thr Val i 1045				

atc aag gaa gac agc ttc ttg cag cga tac agc tca gac ccc aca ggc Ile Lys Glu Asp Ser Phe Leu Gln Arg Tyr Ser Ser Asp Pro Thr Gly 1060 1065 1070 1075
gcc ttg act gag gac agc ata gac gac acc ttc ctc cca gtg cct gaa 3439 Ala Leu Thr Glu Asp Ser Ile Asp Asp Thr Phe Leu Pro Val Pro Glu 1080 1085 1090
tac ata aac cag tcc gtt ccc aaa agg ccc gct ggc tct gtg cag aat Tyr Ile Asn Gln Ser Val Pro Lys Arg Pro Ala Gly Ser Val Gln Asn 1095 1100 1105
cct gtc tat cac aat cag cct ctg aac ccc gcg ccc agc aga gac cca 3535 Pro Val Tyr His Asn Gln Pro Leu Asn Pro Ala Pro Ser Arg Asp Pro 1110 1115 1120
cac tac cag gac ccc cac agc act gca gtg ggc aac ccc gag tat ctc 3583 His Tyr Gln Asp Pro His Ser Thr Ala Val Gly Asn Pro Glu Tyr Leu 1125 1130 1135
aac act gtc cag ccc acc tgt gtc aac agc aca ttc gac agc cct gcc 3631 Asn Thr Val Gln Pro Thr Cys Val Asn Ser Thr Phe Asp Ser Pro Ala 1140 1145 1150 1155
cac tgg gcc cag aaa ggc agc cac caa att agc ctg gac aac cct gac His Trp Ala Gln Lys Gly Ser His Gln Ile Ser Leu Asp Asn Pro Asp 1160 1165 1170
tac cag cag gac ttc ttt ccc aag gaa gcc aag cca aat ggc atc ttt 3727 Tyr Gln Gln Asp Phe Phe Pro Lys Glu Ala Lys Pro Asn Gly Ile Phe 1175 1180 1185
aag ggc tcc aca gct gaa aat gca gaa tac cta agg gtc gcg cca caa 3775 Lys Gly Ser Thr Ala Glu Asn Ala Glu Tyr Leu Arg Val Ala Pro Gln 1190 1195 1200
agc agt gaa ttt att gga gca tga ccacggagga tagtatgagc cctaaaaatc 3829 Ser Ser Glu Phe Ile Gly Ala 1205 1210
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Leu Glu Ile Thr Tyr Val Gln Arq Asn Tyr Asp Leu Ser Phe Leu Lys
                                         75
Thr Ile Gln Glu Val Ala Gly Tyr Val Leu Ile Ala Leu Asn Thr Val
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Glu Arg Ile Pro Leu Glu Asn Leu Gln Ile Ile Arg Gly Asn Met Tyr
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Tyr Glu Asn Ser Tyr Ala Leu Ala Val Leu Ser Asn Tyr Asp Ala Asn
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                                               125
Lys Thr Gly Leu Lys Glu Leu Pro Met Arg Asn Leu Gln Glu Ile Leu
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                                            140
His Gly Ala Val Arg Phe Ser Asn Asn Pro Ala Leu Cys Asn Val Glu
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                                       155
Ser Ile Gln Trp Arg Asp Ile Val Ser Ser Asp Phe Leu Ser Asn Met
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Ser Met Asp Phe Gln Asn His Leu Gly Ser Cys Gln Lys Cys Asp Pro
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Ser Cys Pro Asn Gly Ser Cys Trp Gly Ala Gly Glu Glu Asn Cys Gln
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Lys Leu Thr Lys Ile Ile Cys Ala Gln Gln Cys Ser Gly Arg Cys Arg
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                                            220
Gly Lys Ser Pro Ser Asp Cys Cys His Asn Gln Cys Ala Ala Gly Cys
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                                        235
Thr Gly Pro Arg Glu Ser Asp Cys Leu Val Cys Arg Lys Phe Arg Asp
               245
                                    250
Glu Ala Thr Cys Lys Asp Thr Cys Pro Pro Leu Met Leu Tyr Asn Pro
           260
                               265
Thr Thr Tyr Gln Met Asp Val Asn Pro Glu Gly Lys Tyr Ser Phe Gly
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                                                285
Ala Thr Cys Val Lys Lys Cys Pro Arg Asn Tyr Val Val Thr Asp His
                        295
Gly Ser Cys Val Arg Ala Cys Gly Ala Asp Ser Tyr Glu Met Glu Glu
                    310
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Asp Gly Val Arg Lys Cys Lys Cys Glu Gly Pro Cys Arg Lys Val
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                                    330
Cys Asn Gly Ile Gly Ile Gly Glu Phe Lys Asp Ser Leu Ser Ile Asn
            340
                                345
Ala Thr Asn Ile Lys His Phe Lys Asn Cys Thr Ser Ile Ser Gly Asp
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                                                365
Leu His Ile Leu Pro Val Ala Phe Arg Gly Asp Ser Phe Thr His Thr
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Pro Pro Leu Asp Pro Gln Glu Leu Asp Ile Leu Lys Thr Val Lys Glu
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                                        395
Ile Thr Gly Phe Leu Leu Ile Gln Ala Trp Pro Glu Asn Arg Thr Asp
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                                    410
Leu His Ala Phe Glu Asn Leu Glu Ile Ile Arg Gly Arg Thr Lys Gln
                                425
His Gly Gln Phe Ser Leu Ala Val Val Ser Leu Asn Ile Thr Ser Leu
                            440
Gly Leu Arg Ser Leu Lys Glu Ile Ser Asp Gly Asp Val Ile Ile Ser
                       455
                                            460
Gly Asn Lys Asn Leu Cys Tyr Ala Asn Thr Ile Asn Trp Lys Lys Leu
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                                       475
Phe Gly Thr Ser Gly Gln Lys Thr Lys Ile Ile Ser Asn Arg Gly Glu
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Asn Ser Cys Lys Ala Thr Gly Gln Val Cys His Ala Leu Cys Ser Pro
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Glu Gly Cys Trp Gly Pro Glu Pro Arg Asp Cys Val Ser Cys Arg Asn
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Val Ser Arg Gly Arg Glu Cys Val Asp Lys Cys Asn Leu Leu Glu Gly
                        535
                                            540
Glu Pro Arg Glu Phe Val Glu Asn Ser Glu Cys Ile Gln Cys His Pro
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                                        555
Glu Cys Leu Pro Gln Ala Met Asn Ile Thr Cys Thr Gly Arg Gly Pro
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                                    570
Asp Asn Cys Ile Gln Cys Ala His Tyr Ile Asp Gly Pro His Cys Val
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                                585
Lys Thr Cys Pro Ala Gly Val Met Gly Glu Asn Asn Thr Leu Val Trp
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                            600
Lys Tyr Ala Asp Ala Gly His Val Cys His Leu Cys His Pro Asn Cys
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Thr Tyr Gly Cys Thr Gly Pro Gly Leu Glu Gly Cys Pro Thr Asn Gly
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Pro Lys Ile Pro Ser Ile Ala Thr Gly Met Val Gly Ala Leu Leu Leu
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Leu Leu Val Val Ala Leu Gly Ile Gly Leu Phe Met Arg Arg Arg His
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Ile Val Arg Lys Arg Thr Leu Arg Arg Leu Leu Gln Glu Arg Glu Leu
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Val Glu Pro Leu Thr Pro Ser Gly Glu Ala Pro Asn Gln Ala Leu Leu
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                                            700
Arg Ile Leu Lys Glu Thr Glu Phe Lys Lys Ile Lys Val Leu Gly Ser
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Gly Ala Phe Gly Thr Val Tyr Lys Gly Leu Trp Ile Pro Glu Gly Glu
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Lys Val Lys Ile Pro Val Ala Ile Lys Glu Leu Arg Glu Ala Thr Ser
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                                745
Pro Lys Ala Asn Lys Glu Ile Leu Asp Glu Ala Tyr Val Met Ala Ser
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Val Asp Asn Pro His Val Cys Arg Leu Leu Gly Ile Cys Leu Thr Ser
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Thr Val Gln Leu Ile Thr Gln Leu Met Pro Phe Gly Cys Leu Leu Asp
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                                        795
Tyr Val Arg Glu His Lys Asp Asn Ile Gly Ser Gln Tyr Leu Leu Asn
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                                    810
Trp Cys Val Gln Ile Ala Lys Gly Met Asn Tyr Leu Glu Asp Arg Arg
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                                825
Leu Val His Arg Asp Leu Ala Ala Arg Asn Val Leu Val Lys Thr Pro
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Gln His Val Lys Ile Thr Asp Phe Gly Leu Ala Lys Leu Leu Gly Ala
                        855
Glu Glu Lys Glu Tyr His Ala Glu Gly Gly Lys Val Pro Ile Lys Trp
                    870
                                        875
Met Ala Leu Glu Ser Ile Leu His Arg Ile Tyr Thr His Gln Ser Asp
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                                    890
Val Trp Ser Tyr Gly Val Thr Val Trp Glu Leu Met Thr Phe Gly Ser
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Lys Pro Tyr Asp Gly Ile Pro Ala Ser Glu Ile Ser Ser Ile Leu Glu
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Lys Gly Glu Arg Leu Pro Gln Pro Pro Ile Cys Thr Ile Asp Val Tyr
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Phe Arg Glu Leu Ile Ile Glu Phe Ser Lys Met Ala Arg Asp Pro Gln
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Arg Tyr Leu Val Ile Gln Gly Asp Glu Arg Met His Leu Pro Ser Pro
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Thr Asp Ser Asn Phe Tyr Arg Ala Leu Met Asp Glu Glu Asp Met Asp
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Asp Val Val Asp Ala Asp Glu Tyr Leu Ile Pro Gln Gln Gly Phe Phe
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Ser Ser Pro Ser Thr Ser Arg Thr Pro Leu Leu Ser Ser Leu Ser Ala
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Thr Ser Asn Asn Ser Thr Val Ala Cys Ile Asp Arg Asn Gly Leu Gln
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              1045
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Ser Cys Pro Ile Lys Glu Asp Ser Phe Leu Gln Arg Tyr Ser Ser Asp
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                             1065
Pro Thr Gly Ala Leu Thr Glu Asp Ser Ile Asp Asp Thr Phe Leu Pro
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                          1080
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Val Pro Glu Tyr Ile Asn Gln Ser Val Pro Lys Arg Pro Ala Gly Ser
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                                          1100
Val Gln Asn Pro Val Tyr His Asn Gln Pro Leu Asn Pro Ala Pro Ser
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Arg Asp Pro His Tyr Gln Asp Pro His Ser Thr Ala Val Gly Asn Pro
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Glu Tyr Leu Asn Thr Val Gln Pro Thr Cys Val Asn Ser Thr Phe Asp
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Ser Pro Ala His Trp Ala Gln Lys Gly Ser His Gln Ile Ser Leu Asp
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Asn Pro Asp Tyr Gln Gln Asp Phe Pro Lys Glu Ala Lys Pro Asn
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Gly Ile Phe Lys Gly Ser Thr Ala Glu Asn Ala Glu Tyr Leu Arg Val
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<223> Murine EGFR

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gca ggt ggg gcg ttg gag gaa aag aaa gtc tgc Ala Gly Gly Ala Leu Glu Glu Lys Lys Val Cys 25 30	
agg ctc acc caa ctg ggc act ttt gaa gac cac Arg Leu Thr Gln Leu Gly Thr Phe Glu Asp His 40 45	
agg atg tac aac aac tgt gaa gtg gtc ctt ggg Arg Met Tyr Asn Asn Cys Glu Val Val Leu Gly 55 60	
tat gtg caa agg aat tac gac ctt tcc ttc tta Tyr Val Gln Arg Asn Tyr Asp Leu Ser Phe Leu 70 75	
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ttg gag aac ctg cag atc atc agg gga aat gct Leu Glu Asn Leu Gln Ile Ile Arg Gly Asn Ala 105	
tat gcc tta gcc atc ctg tcc aac tat ggg aca Tyr Ala Leu Ala Ile Leu Ser Asn Tyr Gly Thi 120	
agg gaa ctg ccc atg cgg aac tta cag gaa atc Arg Glu Leu Pro Met Arg Asn Leu Gln Glu Ile 135	
cga ttc agc aac aac ccc atc ctc tgc aat atc Arg Phe Ser Asn Asn Pro Ile Leu Cys Asn Met 150	
agg gac atc gtc caa aac gtc ttt atg agc aac Arg Asp Ile Val Gln Asn Val Phe Met Ser Ass 165 170 175	n Met Ser Met Asp Leu
cag agc cat ccg agc agt tgc ccc aaa tgt gat Gln Ser His Pro Ser Ser Cys Pro Lys Cys Asp 185 190	
gga agc tgc tgg gga gga gga gag gag aac tgc Gly Ser Cys Trp Gly Gly Glu Glu Asn Cys 200 205	
atc atc tgt gcc cag caa tgt tcc cat cgc tgt	t cgt ggc agg tcc ccc 907

Ile	Ile	Cys 215	Ala	Gln	Gln	Cys	Ser 220	His	Arg	Cys	Arg	Gly 225	Arg	Ser	Pro	
					aac Asn											955
	-	-	_	-	gtc Val 250	-		-			-		_		_	1003
					cca Pro											1051
_	_	-			gaa Glu		_		_			_		-		1099
_	_	_		_	aac Asn					-				_	_	1147
_	_	_			gac Asp			_		-	-	-			_	1195
					gat Asp 330			_	_		-	_				1243
			-		aaa Lys	_						-				1291
					tgc Cys		-		_		-				_	1339
					ggg Gly											1387
					att Ile											1435
					tgg Trp 410											1483
					ata Ile											1531
					ggc Gly											1579

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	acc aaa atc at Thr Lys Ile Me 490	_		-	
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Gly Pro Glu	ccc agg gac to Pro Arg Asp Cy 520				
	gtg gag aaa to Val Glu Lys Cy			-	
	aat tot gaa to Asn Ser Glu Cy 55	ys Ile Gln	-		
	aac atc acc to Asn Ile Thr Cy 570				
	cac tac att ga His Tyr Ile As 585				
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					gga Gly											2347
					ttc Phe											2395
					aag Lys 730											2443
					atc Ile											2491
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			_	_	cgc Arg		-			_	_					2587
-				_	ctc Leu	_				_		_	-			2635
					aac Asn 810											2683
	_		_	_	ggc Gly	_			_	-	-		-	_		2731
	_	_	_	_	gcc Ala			_								2779
_	_			-	ttt Phe		_	-		_			_	-		2827
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_	-				cac His 890	-						-	-	_		2923

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					gca Ala											3019
_	_			_	cca Pro			-			_	-				3067
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					ttc Phe 970											3163
	-		_		gat Asp	-	_	_		_		_			_	3211
		Phe		_	gcc Ala	_	Met	_	_		_	Met		-	-	3259
-	Asp	_	-	-	tat Tyr	Leu			_		Ğĺy				-	3307
Pro		-	_		act Thr			_	-	Ser	_	_	-		-	3355
	Asn			Val	gct Ala 1050	_			Arg			-	_	Arg	-	3403
			Ala		ttg Leu			Tyr					Thr			3451
		Glu	-		ata Ile	_	Āsp	_				Val		-		3499
	Asn				ccc Pro	Lys					Ser					3547
Val				_	ccc Pro	_			-	Pro		-	-	_		3595
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act gcc cag cct acc tgt ctc agt agt ggg ttt aac agc cct gca ctc Thr Ala Gln Pro Thr Cys Leu Ser Ser Gly Phe Asn Ser Pro Ala Leu 1145 1150 1155	3691
tgg atc cag aaa ggc agt cac caa atg agc cta gac aac cct gac tac Trp Ile Gln Lys Gly Ser His Gln Met Ser Leu Asp Asn Pro Asp Tyr 1160 . 1165 1170	3739
cag cag gac ttc ttc ccc aag gaa acc aag cca aat ggc ata ttt aag Gln Gln Asp Phe Phe Pro Lys Glu Thr Lys Pro Asn Gly Ile Phe Lys 1175 1180 1185	3787
ggc ccc aca gct gaa aat gca gag tac cta cgg gtg gca cct cca agc Gly Pro Thr Ala Glu Asn Ala Glu Tyr Leu Arg Val Ala Pro Pro Ser 1190 1195 1200	3835
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<220>

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<223> Murine EGFR

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 Gly
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 Lys
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 Val
 Cys
 Gln

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 Thr
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 Gly
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 Phe
 Glu
 Asp
 His
 Phe

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 Ser
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 Glu
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 Gly
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 Ser
 Phe
 Leu
 Lys

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 75
 80

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Tyr Glu Asn Thr Tyr Ala Leu Ala Ile Leu Ser Asn Tyr Gly Thr Asn
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Arg Thr Gly Leu Arg Glu Leu Pro Met Arg Asn Leu Gln Glu Ile Leu
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Ile Gly Ala Val Arg Phe Ser Asn Asn Pro Ile Leu Cys Asn Met Asp
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Lys Leu Thr Lys Ile Ile Cys Ala Gln Gln Cys Ser His Arg Cys Arg
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Thr Gly Pro Arg Glu Ser Asp Cys Leu Val Cys Gln Lys Phe Gln Asp
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Thr Thr Tyr Gln Met Asp Val Asn Pro Glu Gly Lys Tyr Ser Phe Gly
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Cys Asn Gly Ile Gly Ile Gly Glu Phe Lys Asp Thr Leu Ser Ile Asn
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Gly Asn Arg Asn Leu Cys Tyr Ala Asn Thr Ile Asn Trp Lys Lys Leu
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Gly Ser Lys Pro Tyr Asp Gly Ile Pro Ala Ser Asp Ile Ser Ser Ile
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Pro Lys Phe Arg Glu Leu Ile Leu Glu Phe Ser Lys Met Ala Arg Asp
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gta gtt tca aaa Val Val Ser Lys					520
agc tgt cct gad Ser Cys Pro Glo 30	Gly Thr Leu			• • •	68
ggt cct gca ccc Gly Pro Ala Pro 45					516
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											cag Gln			1192
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											aag Lys			1288
							_				gaa Glu	_	_	 1336
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											gca Ala			1480
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		tgt Cys														1864
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		cag Gln														2008
-		aag Lys 525				_			_	_	_				_	2056
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		cca Pro	_		_	_								_	_	2344
		gcc Ala														2392
		act Thr														2440
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Asn Ser Asp Ser (cc cac gat ggg tac to er His Asp Gly Tyr Cy 980	
	Met Tyr Ile Glu A	gca ttg gac aag tat go ala Leu Asp Lys Tyr Al 195 100	a Cys Asn
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	Tyr Glu Glu Ser S	ngc aga gat gtg agg ag Ser Arg Asp Val Arg Se 175 108	r Arg Arg
		cc tct tgc cct caa co er Ser Cys Pro Gln Pr 1095	
		etc aag aat ggg ggt ca eu Lys Asn Gly Gly Gl 1110	
		at ggg tca atg caa co sp Gly Ser Met Gln Pr 1125	

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tgg att cca gta tcc agt gat aag ggc tcc tgt ccc cag gta atg gag 3928 Trp Ile Pro Val Ser Ser Asp Lys Gly Ser Cys Pro Gln Val Met Glu 1150 1155 1160	3
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	_	_			-	_	_	aga Arg	_						2036
								tgg Trp 570							2084
								gga Gly							2132
		_				_		aga Arg		_					2180
								ctg Leu							2228

											tcc Ser					2276
											atc Ile					2324
											agg Arg					2372
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					Pro						aac Asn					2516
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											gat Asp					2612
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				-	_				_	_	tgg Trp	_			-	2708
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		_				_	-				cag Gln	-	-	-		2804
											gct Ala					2852
											ggg Gly 845					2900
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His A 850	Ala	Arg	Cys	Val	Ser 855	Asp	Gly	Glu	Thr	Ala 860	Glu	Cys	Gln	Cys	Leu 865	
aaa g Lys G																2996
gtg c Val L	_	-	-	_	_	-										3044
act g Thr G					-	_	_	_		-	-					3092
ggg a Gly I			_		_		-		-	_						3140
tgc g Cys A 930																3188
acc t Thr C	_	_		_			_				_	-				3236
acc g Thr A							-	_								3284
agt t Ser T																3332
ggc g Gly V					Ile					Ser						3380
gtt a Val I 1010				Ser					Gln					Arg		3428
tgg g Trp G		_	Arg		_			Gly	_	-		_	Ile	_		3476
gtg g Val A		Val					Leu					Leu				3524
tgg g Trp G	Sly					Arg					Leu					3572
aag a Lys A																3620

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gat ggt acg aat ggt gca gta gta gat gct ggc ctg tct ccc tcc ctg Asp Gly Thr Asn Gly Ala Val Val Asp Ala Gly Leu Ser Pro Ser Leu 1125 1130 1135	3764
cag ctc ggg tca gtg cat ctg act tca tgg aga cag aag ccc cac ata Gln Leu Gly Ser Val His Leu Thr Ser Trp Arg Gln Lys Pro His Ile 1140 1145 1150	3812
gat gga atg ggc aca ggg caa agc tgc tgg att cca cca tca agt gac Asp Gly Met Gly Thr Gly Gln Ser Cys Trp Ile Pro Pro Ser Ser Asp 1155 1160 1165	3860
aga gga ccc cag gaa ata gag gga aac tcc cac cta ccc tcc tac aga Arg Gly Pro Gln Glu Ile Glu Gly Asn Ser His Leu Pro Ser Tyr Arg 1170 1175 1180 1185	3908
cct gtg ggg ccg gag aag ctg cat tct ctc cag tca gct aat gga tcg Pro Val Gly Pro Glu Lys Leu His Ser Leu Gln Ser Ala Asn Gly Ser 1190 1195 1200	3956
tgt cac gaa agg gct cca gac ctg cca cgg cag aca gag cca gtt aag Cys His Glu Arg Ala Pro Asp. Leu Pro Arg Gln Thr Glu Pro Val Lys 1205 1210 1215	4004
tag aaactgggag tagacagaag gtacagaagg gaaaataaca aaccaggctg	4057
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agggtcctga agatagctgc acagctgcag agctgcacag cgggatagct gcgacttttg	4177
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Asn Cys Gln Pro Gly Pro Leu Glu Arg Ser Glu Arg Ser Gly Thr Cys
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Ala Gly Pro Ala Pro Phe Leu Val Phe Ser Gln Gly Lys Ser Ile Ser
Arg Ile Asp Pro Asp Gly Thr Asn His Gln Gln Leu Val Val Asp Ala
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Gly Ile Ser Ala Asp Met Asp Ile His Tyr Lys Lys Glu Arg Leu Tyr
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Trp Val Asp Val Glu Arg Gln Val Leu Leu Arg Val Phe Leu Asn Gly
           100
                                105
Thr Gly Leu Glu Lys Val Cys Asn Val Glu Arg Lys Val Ser Gly Leu
                            120
Ala Ile Asp Trp Ile Asp Asp Glu Val Leu Trp Val Asp Gln Gln Asn
                        135
                                            140
Gly Val Ile Thr Val Thr Asp Met Thr Gly Lys Asn Ser Arg Val Leu
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                                        155
Leu Ser Ser Leu Lys His Pro Ser Asn Ile Ala Val Asp Pro Ile Glu
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                                    170
Arg Leu Met Phe Trp Ser Ser Glu Val Thr Gly Ser Leu His Arg Ala
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                                                    190
His Leu Lys Gly Val Asp Val Lys Thr Leu Leu Glu Thr Gly Gly Ile
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Ser Val Leu Thr Leu Asp Val Leu Asp Lys Arg Leu Phe Trp Val Gln
                        215
                                            220
Asp Ser Gly Glu Gly Ser His Ala Tyr Ile His Ser Cys Asp Tyr Glu
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                                        235
Gly Gly Ser Val Arg Leu Ile Arg His Gln Ala Arg His Ser Leu Ser
                245
                                    250
Ser Met Ala Phe Phe Gly Asp Arg Ile Phe Tyr Ser Val Leu Lys Ser
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Lys Ala Ile Trp Ile Ala Asn Lys His Thr Gly Lys Asp Thr Val Arg
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                                                285
Ile Asn Leu His Pro Ser Phe Val Thr Pro Gly Lys Leu Met Val Val
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                                            300
His Pro Arg Ala Gln Pro Arg Thr Glu Asp Ala Ala Lys Asp Pro Asp
                    310
                                        315
Pro Glu Leu Lys Gln Arg Gly Arg Pro Cys Arg Phe Gly Leu Cys
                325
                                    330
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Glu Arg Asp Pro Lys Ser His Ser Ser Ala Cys Ala Glu Gly Tyr Thr
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Leu Ser Arg Asp Arg Lys Tyr Cys Glu Asp Val Asn Glu Cys Ala Thr
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Gln Asn His Gly Cys Thr Leu Gly Cys Glu Asn Thr Pro Gly Ser Tyr
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His Cys Thr Cys Pro Thr Gly Phe Val Leu Leu Pro Asp Gly Lys Gln
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Cys His Glu Leu Val Ser Cys Pro Gly Asn Val Ser Lys Cys Ser His
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Gly Cys Val Leu Thr Ser Asp Gly Pro Arg Cys Ile Cys Pro Ala Gly
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Ser Val Leu Gly Arg Asp Gly Lys Thr Cys Thr Gly Cys Ser Ser Pro
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Asp Asn Gly Gly Cys Ser Gln Ile Cys Leu Pro Leu Arg Pro Gly Ser
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Trp Glu Cys Asp Cys Phe Pro Gly Tyr Asp Leu Gln Ser Asp Arg Lys
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Ser Cys Ala Ala Ser Gly Pro Gln Pro Leu Leu Phe Ala Asn Ser
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Gln Asp Ile Arg His Met His Phe Asp Gly Thr Asp Tyr Lys Val Leu
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Leu Ser Arg Gln Met Gly Met Val Phe Ala Leu Asp Tyr Asp Pro Val
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                                                525
Glu Ser Lys Ile Tyr Phe Ala Gln Thr Ala Leu Lys Trp Ile Glu Arg
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Ala Asn Met Asp Gly Ser Gln Arg Glu Arg Leu Ile Thr Glu Gly Val
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Asp Thr Leu Glu Gly Leu Ala Leu Asp Trp Ile Gly Arg Arg Ile Tyr
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Trp Thr Asp Ser Gly Lys Ser Val Val Gly Gly Ser Asp Leu Ser Gly
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Lys His His Arg Ile Ile Ile Gln Glu Arg Ile Ser Arg Pro Arg Gly
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Met Ser Pro Arg Ile Glu Ser Ala Ser Leu Gln Gly Ser Asp Arg Val
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Leu Ile Ala Ser Ser Asn Leu Leu Glu Pro Ser Gly Ile Thr Ile Asp
                                    650
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Tyr Leu Thr Asp Thr Leu Tyr Trp Cys Asp Thr Lys Arg Ser Val Ile
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Glu Met Ala Asn Leu Asp Gly Ser Lys Arg Arg Arg Leu Ile Gln Asn
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Asp Val Gly His Pro Phe Ser Leu Ala Val Phe Glu Asp His Leu Trp
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                                            700
Val Ser Asp Trp Ala Ile Pro Ser Val Ile Arg Val Asn Lys Arg Thr
                    710
                                        715
Gly Gln Asn Arg Val Arg Leu Gln Gly Ser Met Leu Lys Pro Ser Ser
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                                    730
Leu Val Val His Pro Leu Ala Lys Pro Gly Ala Asp Pro Cys Leu
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Tyr Arg Asn Gly Gly Cys Glu His Ile Cys Gln Glu Ser Leu Gly Thr
                            760
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Ala Arg Cys Leu Cys Arg Glu Gly Phe Val Lys Ala Trp Asp Gly Lys
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Met Cys Leu Pro Gln Asp Tyr Pro Ile Leu Ser Gly Glu Asn Ala Asp
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Val Ser Gly Met Asn Tyr Glu Asp Asp Cys Gly Pro Gly Gly Cys Gly
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Ser His Ala Arg Cys Val Ser Asp Gly Glu Thr Ala Glu Cys Gln Cys
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                                     875
Cys Val Leu Ala Arg Ser Asp Cys Pro Ser Thr Ser Ser Arg Cys Ile
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                                 890
Asn Thr Glu Gly Gly Tyr Val Cys Arg Cys Ser Glu Gly Tyr Glu Gly
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Asp Gly Ile Ser Cys Phe Asp Ile Asp Glu Cys Gln Arg Gly Ala His
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Asn Cys Ala Glu Asn Ala Ala Cys Thr Asn Thr Glu Gly Gly Tyr Asn
                                         940
                      935
Cys Thr Cys Ala Gly Arg Pro Ser Ser Pro Gly Arg Ser Cys Pro Asp
                                     955
Ser Thr Ala Pro Ser Leu Leu Gly Glu Asp Gly His His Leu Asp Arg
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Asn Ser Tyr Pro Gly Cys Pro Ser Ser Tyr Asp Gly Tyr Cys Leu Asn
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Gly Gly Val Cys Met His Ile Glu Ser Leu Asp Ser Tyr Thr Cys Asn
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Cys Val Ile Gly Tyr Ser Gly Asp Arg Cys Gln Thr Arg Asp Leu Arg
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Trp Trp Glu Leu Arg His Ala Gly Tyr Gly Gln Lys His Asp Ile Met
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Val Val Ala Val Cys Met Val Ala Leu Val Leu Leu Leu Leu Gly
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Met Trp Gly Thr Tyr Tyr Arg Thr Arg Lys Gln Leu Ser Asn Pro
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Pro Lys Asn Pro Cys Asp Glu Pro Ser Gly Ser Val Ser Ser Ser Gly
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Pro Asp Ser Ser Ser Gly Ala Ala Val Ala Ser Cys Pro Gln Pro Trp
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Phe Val Val Leu Glu Lys His Gln Asp Pro Lys Asn Gly Ser Leu Pro
1105 1110 1115
Ala Asp Gly Thr Asn Gly Ala Val Asp Ala Gly Leu Ser Pro Ser
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Leu Gln Leu Gly Ser Val His Leu Thr Ser Trp Arg Gln Lys Pro His
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                            1145
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Asp Arg Gly Pro Gln Glu Ile Glu Gly Asn Ser His Leu Pro Ser Tyr
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Arg Pro Val Gly Pro Glu Lys Leu His Ser Leu Gln Ser Ala Asn Gly
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Leu Ala Leu Phe Ala Leu Gly Ile Val Leu Ala Ala Cys Gln Ala Leu
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                              15
                                                   20
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gag aac agc acg tcc ccg ctg agt gca gac ccg ccc gtg gct gca gca
Glu Asn Ser Thr Ser Pro Leu Ser Ala Asp Pro Pro Val Ala Ala Ala
                                                                    196
gtg gtg tcc cat ttt aat gac tgc cca gat tcc cac act cag ttc tgc
Val Val Ser His Phe Asn Asp Cys Pro Asp Ser His Thr Gln Phe Cys
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ttc cat gga acc tgc agg ttt ttg gtg cag gag gac aag cca gca tgt
Phe His Gly Thr Cys Arg Phe Leu Val Gln Glu Asp Lys Pro Ala Cys
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Val Cys His Ser Gly Tyr Val Gly Ala Arg Cys Glu His Ala Asp Leu
ctg gcc gtg gtg gct gcc agc cag aag aag cag gcc atc acc gcc ttg
                                                                    340
Leu Ala Val Val Ala Ala Ser Gln Lys Lys Gln Ala Ile Thr Ala Leu
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                              95
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gtg gtg gtc tcc atc gtg gcc ctg gct gtc ctt atc atc aca tgt gtg
Val Val Ser Ile Val Ala Leu Ala Val Leu Ile Ile Thr Cys Val
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                         110
                                              115
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ctg ata cac tgc tgc cag gtc cga aaa cac tgt gag tgg tgc cgg gcc
Leu Ile His Cys Cys Gln Val Arg Lys His Cys Glu Trp Cys Arg Ala
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ctc atc tgc cgg cac gag aag ccc agc gcc ctc ctg aag gga aga acc
Leu Ile Cys Arg His Glu Lys Pro Ser Ala Leu Leu Lys Gly Arg Thr
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Ala Cys Cys His Ser Glu Thr Val Val
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Asp Ser His Thr Gln Phe Cys Phe His Gly Thr Cys Arg Phe Leu Val
                         55
Gln Glu Asp Lys Pro Ala Cys Val Cys His Ser Gly Tyr Val Gly Ala
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                                          75
Arg Cys Glu His Ala Asp Leu Leu Ala Val Val Ala Ala Ser Gln Lys
                                      90
Lys Gln Ala Ile Thr Ala Leu Val Val Ser Ile Val Ala Leu Ala
            100
Val Leu Ile Ile Thr Cys Val Leu Ile His Cys Cys Gln Val Arg Lys
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                            120
                                                 125
His Cys Glu Trp Cys Arg Ala Leu Ile Cys Arg His Glu Lys Pro Ser
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Ala Leu Leu Lys Gly Arg Thr Ala Cys Cys His Ser Glu Thr Val Val
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<211> 3776

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<213> Mus musculus

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<220>

<221> misc_feature

<223> Murine TGF-alpha

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aac agc aca tcc ccc ctg agt gac tca ccc gtg gcg gct gca gtg gtg Asn Ser Thr Ser Pro Leu Ser Asp Ser Pro Val Ala Ala Ala Val Val 25 30 35 40	10
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gga acc tgc cgg ttt ttg gtg cag gaa gag aag cca gca tgt gtc tgc 30 Gly Thr Cys Arg Phe Leu Val Gln Glu Glu Lys Pro Ala Cys Val Cys 60 . 65 70	06
cac tot ggg tac gtg ggt gtt cgc tgt gag cat gca gac ctc ctg gct His Ser Gly Tyr Val Gly Val Arg Cys Glu His Ala Asp Leu Leu Ala 75 80 85	54
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Gly Leu Arg Thr Leu Met Leu Arg Asn Asn Arg Ile Ser Cys Ile His 50 55 60

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Xaa Xaa Xaa